## O'Bryen, Barbara

From:

Switzer, Juliet

Sent:

Thursday, October 14, 2004 2:10 PM

To: Subject: O'Bryen, Barbara RE: alignment

sorry. seq id no 58 of serial number 10077176. thanks.

----Original Message----From:

O'Bryen, Barbara

Sent:

Thursday, October 14, 2004 2:10 PM

To:

Switzer, Juliet

Subject:

RE: alignment

It would be easier for me to do this if I had & seq id # & serial number.

----Original Message----

From: Switzer, Juliet

Sent:

Thursday, October 14, 2004 2:02 PM

To: O'Bryen, Barbara

Subject: alignment

HI barb.

Will you please align nucleotides 252-1430 against GenBank X60012? Will you let me know when you've done it- I'll come pick up the print out. Thanks.

Juliet

Lipis Page Blank (uspto)

Run on:

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/translation="MEEPQSDESVEPPLSQETFSDLWKLLPENNVLSPLPSQAMDDLM LSPDIEQWFTEDPGPDEAPRMPEAAPTPAAPAPAAPAPAPAPAPABDLM LSPDDIEQWFTEDPGPDEAPRMPEAAPTRVAAPAAPTPAAAPAPAPAPAPAPAPAPARTSQSVETTYGSQUMTUSTSVETVETVPERCYDLWTVSTPPFGTRVRAM ALYKQSQHMTEVVRRCPHHERCSDSDGLAPPQHLIRVEGULRVEYLDDRNTFRHSVVV PYEPPBYGSDCTTIHYNMCMSSCMGGAMRRPILTITLEDSSGNLLGRNSFEVRVCA CPGRDRRTEEENLRKKGEPHHELPPGSTKRALDNNTSSSPQPKKKPLDGEYFTLQIRG RERFEMENEALELKDAQAGKEPGGSRAHSSHLKSKKGGSTSRHKKLMFKTEGPDS
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/product="p53 transformation suppressor"
                                                        cell line'
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                          /mol type="manny"
/isolate="Burkitts lymphoma
/db xref="taxon:9606"
/cell line="laRC/BL2"
                                                                                                                                                                                                                                                                                     /protein_id="CAA42627.1"
|db_xref="G1:506437"
|db_xref="SWISS-PROT:P04637"
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100.0%; Pred. No. 0;
cive 0; Mismatches
                  'organism="Homo sapiens
                                                                                                                                                                                                  evidence=experimental
                                                                                                                        /gene="p53"
1. .1179
/gene="p53"
/note="cDNA"
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/gene="p53"
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo
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Location/Qualifiers
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Farrell, P.J., Allan, G.J., Shanahan, F., Vousden, K.H. and Crook, T. p53 is frequently mutated in Burkitt's lymphoma cell lines EMBO J. 10 (10), 2879-2887 (1991)
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TOIG of: x60012
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                                                                                                             ; Search time 1 Seconds (without alignments) 2.780 Million cell updates/sec
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X60012
X60012.1 GI:506436
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       GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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x60012
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Homo sapiens (human)
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Farrell, P.J.
Direct Submission
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1 (bases 1 to 1179)

2 rariell, P.J., Allan, G.J., Shanahan, F., Vousden, K.H. and Crook, T. p53 is frequently mutared in Burkitt's lymphoma cell lines

EMBO J. 10 (10), 2879-2887 (1991)
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                                         660
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Submitted (03-JUN-1991) P.J. Farrell, Ludwig Inst for Cancer
St Mary's Hospital Med School, Norfolk Place, London W2 1PG,
mutated p53 transformation suppressor gene.
                                                                                                                                                                                                                                                                                                                               901 CCAGGGAGCACTAAGCGAGCACTGCCCAACAACACCAGCTCCTCTCCCCAGCCAAAGAAG
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TTGCGTGTGGAGTATTTGGATGACAGAAACACTTTTCGACATAGTGTGGTGGTGCCCTAT
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Human mRNA for mutated p53 transformation suppressor gene
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/wol_type="mRNA"
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Homo sapiens (human)
Homo sapiens
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Farrell, P.J.
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X60012.1 GI:506436
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TITLE
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51.2%; Pred. No. 0;
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GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
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- nucleic search, using sw model OM nucleic

September 28, 2004, 12:05:54 ; Search time 16 Seconds (without alignments) 2.637 Million cell updates/sec

Run on:

SEQ58-252-1430 1179

1 atggaggagccgcagtcaga......cagaagggcctgactcagac 1179 Perfect score: Sequence:

IDENTITY NUC Gapox 10.5 Scoring table:

13 seqs, 17894 residues Searched:

Total number of hits satisfying chosen parameters:

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Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

issdb:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## STIMMARTES

Score Match Length DB ID    179   100.0   1182   1 US-08-184-009-215   Sequence 215, 179   100.0   1182   1 US-08-456-215   Sequence 215, 179   100.0   1182   1 US-08-456-215   Sequence 215, 179   100.0   1182   1 US-08-456-736-215   Sequence 215, 179   100.0   1182   1 US-08-460-736-215   Sequence 215, 179   100.0   1182   1 US-08-450-736-215   Sequence 215, 179   100.0   1484   1 US-08-450-736-215   Sequence 215, 179   100.0   1484   1 US-08-450-736-215   Sequence 103, 179   100.0   1512   1 US-08-450-736-215   Sequence 103, 179   100.0   1512   1 US-08-460-736-99   Sequence 99, 179   100.0   1512   1 US-08-450-99   Sequence 99, 179   100.0   1512   1 US-08-450-99   Sequence 215, 26.6   2.3   1182   1 US-08-450-756-215   Sequence 215, 26.6   2.3   1182   1 US-08-450-756-215   Sequence 215, 26.6   2.3   1182   1 US-08-450-756-215   Sequence 215, 26.6   2.3   1182   1 US-08-450-756-103   Sequence 103, 26.6   2.3   1484   1 US-08-450-736-103   Sequence 103, 26.6   2.3   1484   1 US-08-450-736-103   Sequence 103, 26.6   2.3   1484   1 US-08-450-736-103   Sequence 99, 26.6   2.3   1512   1 US-08-450-736-99			400	4 4	ָלְינֶע הַלְינֶע	Trick to	בן קליני ני	4	4 4	44.4	7,4	קקנע	1,44	וממע	ייקקעע	TAME	4	dd v	Tddw	App	App	App	App	App	App	Appl	1	1444	App1
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## ALIGNMENTS

RESULT 1 US-08-184-009-215

61 GACCTATGGAAACTACTTCCTGAAAACAACGTTCTGTCCCCCTTGCCGTCCCAAGCAATG 120 61 GACCTATGGAAACTACTTCTGAAAAAAAAGTTCTGTCCCCTTGCCGTCCCAAGCAATG 120 240 121 GATGATTTGATGCTGTCCCCGGACGATATTGAACAATGGTTCACTGAAGACCCAGGTCCA 180 300 301 AAAACCTACCAGGGCAGCTACGGTTTCCGTCTGGGCTTCTTGCATTCTGGGACAGCCAAG 360 9 TCTGTGACTTGCACGTACTCCCCTCCACCAACAAGATGTTTTGCCAACTGGCCAAGACC 420 361 TCTGTGACCTACCTACTCCCCTGCCCTCAACATGTTTTGCCAACTGGCCAAGAGACC 420 1 ATGGAGGAGCCGCAGTCAGATCCTAGCGTCGAGCCCCCTCTGAGTCAGGAAACATTTTCA 181 GATGAAGCTCCCAGAATGCCAGAGGCTGCTCCCCGCGTGGCCCCTGCACCAGCAGCTCCT 241 ACACCGGCCCTGCACCAGCCCCCTCCTGGCCCCTGTCATCTTCTGTCCTTCCCAG 301 AAAACCTACCAGGGGAGCTACGGTTTCCGTCTGGGCTTCTTGCATTCTGGGACAGGCCAAG Gaps 100.0%; Score 1179; DB 1; Length 1182; 100.0%; Pred. No. 0.00062; tive 0; Mismatches 0; Indels 0; Sequence 215, Application US/08184009
Patent No. 5833975
BERENAL INFORMATION:
APPLICANT: Paoletti, Enzo
APPLICANT: Tartaglia, James
APPLICANT: Tortaglia, James
APPLICANT: Cov. William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
ADDRESSED: Curtis, Movenue
STREET: 530 Fifth Avenue
CITY: New York
STRATE: NY
COUNTRY: USA ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/184,009
FILING DATE: 19-JAN-1994
CLASSIFICATION: 4355
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S. 454310-2530 NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 45431
REFERENCE/DOCKET NUMBER: 45431
TELECOMMUNICATION INFORMATION:
TELEFAX: (212) 840-333
TELEFAX: (212) 840-373
TELEX: 425066CURING
TELEX: 42506CURING
TELEX: 1208 BAD NO: 215:
SEQUENCE CHARACTERISTICS:
LENGTH: 1182 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear Query Match 100. Best Local Similarity 100. Matches 1179; Conservative ; MOLECULE TYPE: CDNA US-08-184-009-215 361 ò 임 à 셤  $\delta$ d d ð ò g à g ਨੇ

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GCCATCTACAAGCAGTCACAGGACATGACGGAGGTTGTGAGGCGCTGCCCCCCCATGAG 540
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      PatentIn Release #1.0, Version #1.25
                                                                                                                                                                            454310-2530
                                 APPLICATION NUMBER: US/08/458,356
FILING DATE: 02-UUN-1995
CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/184,009
FILING DATE: 19-UAN-1994
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
                                                                                                                                           NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 45.
TELECOMMUNICATION: TELECOMMUNICATION: (212) 840-3333
                                                                                                                                                                                                                                     TELEX: 425066CURTMS
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 1182 base pairs
                                                                                                                                                                                                                       (212) 840-0712
                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                          TELEPHONE:
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TGCCCTGTGCAGCTGTGGTTGATTCCACACCCCCGCCCGGCACCCGCGTCCGCGCCATG
               TIGCGIGIGGAGIATITIGGAIGACAGAAACACTITITCGACATAGIGIGGIGGIGCCCIAI
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APPLICANT: Tartaglia, James
APPLICANT: Cox, William I.
TILE OF INVENTION: RECOMBINANT VIRUS
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 215, Application US/08458356 Patent No. 5942235 GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
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                                                        AGTGGTAATCTACTGGGACGGAACAGCTTTGAGGTGCTGTTTGTGCCTGTCCTGGGAGA
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APPLICANT: PEPSTEIN, STEPHEN E.
APPLICANT: SPEIR, EDITH
APPLICANT: ZHOU, YI FU
APPLICANT: ZHOU, YI FU
APPLICANT: ZHOU, YI FU
APPLICANT: BUILE, LORNE
APPLICANT: PINCUS, STEVEN
TITLE OF INVENTION: RESTENOSIS/ATHEROSCLEROSIS DIAGNOSIS,
TITLE OF INVENTION: PROPHYLAXIS AND THERAPY
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: CURTIS, MORRIS & SAFFORD, P.C.
STEERT: SAO FIFTH AVENUE
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPREATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: US/08/796,101
FILING DATE: 05-FEB-1997
CLASSIFICATION: 424
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US-08-796-101-46
Sequence 46, Application US/08796101
Patent No. 6183752
GENERAL INFORMATION:
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ATTORNEY/AGENT INPORMATION:
NAME: KOMALKI, THOMAS J.
REGISTRATION NUMBER: 32,147
TELECOMMUNICATION INFORMATION:
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STATE: NY
COUNTRY:
ZIP: 1003
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                                                                                                                                                                                                     Length 1182
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                                                                                                                                                                                              100.0%; Score 1179; DB 1;
100.0%; Pred. No. 0.00062;
tive 0; Mismatches 0;
           TELEFAX: (212) 764-5574

INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 1182 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
NOLECULE TYPE: DNA (genomic)
US-08-796-101-46
(212) 840-3333
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APPLICANT: Tartaqlia, James
APPLICANT: Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
STREET: 530 Fifth Avenue
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                            AAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC 1179
                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,736
FILING DATE: 02-UNN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/184,009
FILING DATE: 19-JAN-1994
ATTORNEY AGENT INFORMATION:
NAMME: Frommer, Milliam S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
TELECOMMUNICATION INFORMATION:
MANTER COMPUNICATION INFORMATION:
MANTER COMPUNICATION INFORMATION:
MANTER COMPUNICATION INFORMATION:
MANTER COMPUNICATION INFORMATION:
MANTER COMPUTER: 0300 03330
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LELEFAX: (212) 840-333
TELEX: 42506CURING
INPORMATION FOR SEQ ID NO: 215:
SEQUENCE CHARACTERISTICS:
LENGTH: 1182 base pairs
TYRE: mucleic acid
STRANDENNES: sind'
MOLECULE
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 215, Application US/08460736 Patent No. 6265189
                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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US-08-460-736-215
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' Patent No. 5833975
' GENERAL INFORMATION'
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US-08-184-009-103
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Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/535,370
FILING DATE: 24-Mar-2000
CLASSIFICATION: <u cutomomono
CLASSIFICATION: <u cutomomono
CLASSIFICATION: <u cctoology

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100.0%; Pred. No. 0.00062;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: «UDKNOWN»
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Curtis, Morris & Safford
STREET: 530 Fifth Avenue
CITY: New York
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                                                                                                                                                                                                                      Sequence 215, Application US/09535370
Patent No. 6537594
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 1182 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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TELEX: 425066CURTMS
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                         APPLICANT: Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Matches 1179; Conservative
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COUNTRY: USA
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US-09-535-370-215
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                  TTGCGTGTGGAGTATTTGGATGACAGAAACACTTTTCGACATAGTGTGGGTGCTGCCTAT
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Patent No. 594235
GENERAL INFORMATION:
APPLICANT: Pacletti, Enzo
APPLICANT: Tartaqlia, James
APPLICANT: Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cuttis, Morris & Safford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/458,356
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COMPUTER READABLE FORM:
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US-08-458-356-103
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                         APPLICANT: Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
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100.0%; Pred. No. 0.00049;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                               ADDRESSEE: Curtis, Morris & Safford STRRET: 530 Fifth Avenue CITY: New York STATE: NY COUNTRY: USA
                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 840-0712
TELEX: 425066CURMS
INFORMATION FOR SEQ ID NO: 103:
SEQUENCE CHARACTERISTICS:
LENGTH: 1484 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 45
TELECOMMUNICATION INFORMATION
TELEPHONE: (212) 840-0712
TELEPAX: (212) 840-0712
          Tartaglia, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 1179; Conservative
                                                                        CORRESPONDENCE ADDRESS: ADDRESSEE: Curtis, Mc
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US-08-184-009-103
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                                                                                                                1073 GACCGGCGCACAGAGGAAGAGAATCTCCGCAAGAAAGGGGGGCCTCACCACGAGCTGCCC
                               TCCTGCATGGGCGGCATGAACCGGAGGCCCATCCTCACCATCACCACGAAGACTCC
                                                                                                                                                          841 GACCGGCGCACAGAGGAAGGAATCTCCGCAAGAAAGGGGAGCCTCACCACGAGCTGCCC
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APPLICANT: Tartaglia, James
APPLICANT: Tartaglia, James
APPLICANT: Cox, Milliam I.
APPLICANT: Cox, Milliam I.
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESSS:
STREET: 530 Fifth Avenue
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEB PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,736
FILING DATE: 02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION INMERS: US 08/184,009
FILING DATE: 19-7AN-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-460-736-103
; Sequence 103, Application US/08460736
; Patent No. 6265189
; GENERAL INFORMATION:
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TELEFAX: (212) 840-712
TELEX: 425066CURTMS
INFORMATION FOR SEQ ID NO: 103;
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100.0%; Pred. No. 0.00049;
Ative 0; Mismatches 0;
FILING DATE: 02-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION DATA:
PLING DATE: 19-JAN-1994
ATTONEV/AGBAY INFORMATION:
NAME: PROMET: William S.
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 454310-2530
TELEFRANCE/DOKKET NUMBER: 454310-2530
TELEFAM: (212) 840-333
TELEFAM: 425066CURTMS
INFORMATION FOR SEQ ID NO: 103:
SEQUENCE CHARACTERISTICS:
LENGTH: 1484 base pairs
TYPE: NUCLEic acid
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Best Local Similarity 100.
Matches 1179; Conservative
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                                                                                   Length 1484;
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; Pred. No. 0.00049;
0; Mismatches 0;
                                                                                  Query Match
Best Local Similarity 100.0%;
Matches 1179; Conservative 0
1484 base pairs
            TYPE: nucleic acid
STRANDEDNESS: single
                                                CDNA
                                     linear
                                   ; TOPOLOGY: lir;
; MOLECULE TYPE:
US-08-460-736-103
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Cox, William I
TILE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
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REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   & Safford
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APPLICATION NUMBER: US/09/535,370
FILING DATE: 24-Mar-2000
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/460,736
                                                         AAACCACTGGATGGAGAATATTTCACCC
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STREET: 530 Fifth Avenue
CITY: New York
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US-09-535-370-103
Sequence 103, Application US/09535370
Patent No. 6537594
GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
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TELEX: 425066CURTMS
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ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: single
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COMPUTER READABLE FORM:
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Oy 1141 AAAAAACTCATGTT:	RESULT 10 US-08-184-009-99 ; Sequence 99, Application US; Patent No. 5833975 ; GENERAL INFORMATION: ; APPLICANT: Paoleri En	APPLICANT: Tartaglia, J APPLICANT: Cox, William TITLE OF INVENTION: REC WUMBER OF SEQUENCES: 21	CORRESPONDENCE ADDRESS:	. 136 136 136 136	COMPUTER: 1BM PC comp ; OPERATING SYSTEM: PC- ; SOFTWARE: PATENTIR RE ; CURRENT APPLICATION DATA	; APPLICATION NUMBER; U ; FILING DATE: 19-AN-1 ; CLASSIFICATION: 435 ; ATTORNEY/AGENT INFORMAT	NAME: Frommer, Willia REGISTRATION NUMBER: REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORM:	; TELEPHONE: (212) 840-7; TELEFAX: (212) 840-07; TELEX: 42506GURINS; INFORMATION FOR SEQ ID NO:	SEQUENCE CHARACTERISTICS  LENGTH: 1512 base pai  TYPE: NUCleic acid  STRANDENESS: sincle		Query Match Best Local Similarity 100 Matches 1179; Conservative	Qy 1 ATGGAGGCCGCAG:	Oy 61 GACCTATGGAAACTAC	2y 121 GATGATTGADGCTGT 	Oy 181 GATGAAGCTCCCAGAA	Qy 241 ACACCGGCGGCCCCTG	Qy 301 AAAACCTACCAGGGCA
	121 GATCATTICATGCTGTCCCCGGACGATATTGAACAATGGTTCACTGAAGACCCAGGTCCA 180		AAAACCTACCAGGGCAGCTACGGTTTCCGTCTGGGCTTCTTGGATTCTGGGACAGCCAAG		421 TGCCCTGTGCAGCTGTGATTCCACACCCCGGCACCGGGACCGGGGCGTG 480 	481 GCCATCTACAAGCAGTCACAGTGACGGAGGTTGTGAGGCGCTGCCCCCACCATGAG 540 	541 CGCTGCTCAGATAGCGATGGTCTGGCCCCTCCTCAGCATCTTATCCGAGTGGAAGGAA		661 GAGCCGCCTGAGGTTGGCTCTGACTGTACCACCATCCACTACAACTACATGTGTAACAGT 720 	721 TCCTGCATGGGCGGCATGAACCGGAGGCCCATCCTCACCATCATCACCTGGAAGACTCC 780 	781 AGTGGTAATCTACTGGGAACGGAACAGCTTTGAGGTGCGTGTTTGTGCCTGTCCTGGGAAA 840 	841 GACCGGCGCACAGAGGAAGAGATCTCCGCAAGAAAGGGGGGGCTCACCACGAGCTGCCC 900 			1021 TTCCGAGAGCTGAATGAGGCCTTGGAACTCAAGGATGCCCAGGCTGGGAAGGAGCCAGGG 1080 	1081 GGGAGCAGCTCACTCCAGCCACCTGAAGTCCAAAAAGGGTCAGTCTACCTCCCGCCAT 1140 	
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ACTICCTGAAAACAACGITCTGTCCCCCTTGCCGTCCCAAGCAAG 120 AATGCCAGAGGCTGCTCCCCGCGTGGCCCCTGCACCAGCAGCTCCT 240 IGCACCAGCCCCTCCTGGCCCCTGTCATCTTCTGTCCCTTCCCAG 300 AAAACCTACCAGGGCAGCTACGGTTTCCGTCTGGGCTTCTTGCATTCTGGGACAGCCAAG 360 0; Gaps 0.0%; Score 1179; DB 1; Length 1512; 0.0%; Pred. No. 0.00048; e 0; Mismatches 0; Indels 0; TCAAGACAGAAGGCCTGACTCAGAC 1179 am I. ECOMBINANT VIRUS IMMUNOTHERAPY 217 disk mpatible C-DOS/MS-DOS Release #1.0, Version #1.25 Morris & Safford venue TA: US/08/184,009 -1994 JS/08184009 ): 99: 38: tirs Enzo James

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IBM PC Compatible

New York USA

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COMPUTER: LIEM P. COUNDALLING
COMPUTER: DELEM P. COUNDALLING
SOFTWARE: PATENTIN F0-005/MS-005
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/458,356
FILING DATE: 02-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: US 08/184,009
FILING DATE: 19-JAN-1994
ATTONNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET WINDER: 25,506
REFERENCE/DOCKET WINDER: 25,506
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFERK: (212) 840-0712
                                                                                                                                                                                                                                                                               TELEX: 425066CURTMS
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 1512 base pairs
                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                  ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: CDNA
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                        COUNTRY:
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                                                  1 ATGGAGGCGCAGTCAGATCCTAGCGTCGAGCCCCTCTGAGTCAGGAAACATTTTCA
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Length 1512;
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100.0%; Score 1179; DB 1;
100.0%; Pred. No. 0.00048;
iive 0; Mismatches 0;
     Query Match 100.
Best Local Similarity 100.
Matches 1179; Conservative
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| Sequence 99, Application US/08458356 | Sequence 99, Application US/08458356 | Patent No. 5942235 | Patent No. 5942235 | GENERAL INPORMATION: APPLICANT: Pacletti, Enzo APPLICANT: Tartaglia, James APPLICANT: Tox, William I. TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY NUMBER OF SEQUENCES: 217 | CORRESPONDENCE ADDRESS: CARESPONDENCE ADDRESS: STREET: 530 Fifth Avenue

RESULT 11 US-08-458-356-99

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61
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            CGCTGCTCAGATAGCGATGGTCTGGCCCCTCCTCAGCATCTTATCCGAGTGGAAAT
                                           TTGCGTGTGGAGTATTTGGATGACAGAAACACTTTTCGACATAGTGTGGTGGTGCCCTAT
                                                                                          GAGCCCCTGAGGTTGGCTCTGACTGTACCACCATCCACTACAACTACATGTGTAAACAGT
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Fatent No. 6265189
GENERAL INFORMATION
APPLICANT: Paoletti, Enzo
APPLICANT: Tartaglia, James
APPLICANT: Tartaglia, James
APPLICANT: Cox, William I.
ITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
STREET: Say Fifth Avenue
CITY: New York
STATE: NY
COMPUTER: NA
ZIP: 10036
ZIP: 10036
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1141 AAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,736
FLING DATE: 02-UN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/184,009
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US-08-460-736-99
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                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 1179; DB 1;
100.0%; Pred. No. 0.00048;
tive 0; Mismatches 0;
                              NAME: Frommer, William S.

REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
TELEPAN: (212) 840-333
TELEPAN: (212) 840-312
TELENGTH: 1512 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
FILING DATE: 19-JAN-1994
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S
                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100.
Matches 1179; Conservative
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MOLECULE TYPE: CDNA
US-08-460-736-99
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                CCAGGGGACACTAAGCGAGCACTGCCCAACAACACCAGCTCCTCTCCCCAGCCAAAGAAG
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AGTGGTAATCTACTGGGACGGAACAGCTTTGAGGTGCGTGTTTGTGCCTGTCCTGGGAGA
                                                      GACCGGCGCACAGAGAGAGATCTCCGCAAGAAAGGGGGAGCCTCACCACGAGCTGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 24-Mar-2000
CLASSIFICATION: «Unknown.»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Pacletti, Enzo
Tartaglia, James
Cox, William I.
TITLE OF INVENTION RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                          1409 AAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC 1447
                                                                                                                                                                                                                                                                                                                                           AAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Curtis, Morris & Safford
STREET: 530 Fifth Avenue
CIIY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/460,736
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 99, Application US/09535370 Patent No. 6537594 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCE CHARACTERISTICS:
LENGTH: 1512 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 840-0712
TELEX: 425066CURTMS
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COMPUTER READABLE FORM:
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                                                       Length 1512;
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                                                      100.0%; Score 1179; DB 1;
100.0%; Pred. No. 0.00048;
Live 0; Mismatches 0;
; MOLECULE TYPE: CDNA; SRQUENCE DESCRIPTION: SEQ ID NO: US-09-535-370-99
                                                                                     Conservative
                                                             Query Match
Best Local Similarity
Matches 1179; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
APPLICANT: Tartaglia, James
APPLICANT: Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

FILING DATE:

APPLICATION TOWNER:

CLASSIPICATION:

ATTORNEY AGENT INFORMATION:

NAME:

REGISTRATION NUMBER: 25,506

REGISTRATION NUMBER: 25,506

REGISTRATION NUMBER: 25,506

REGISTRATION NUMBER: 25,506

REGISTRATION NUMBER: 454310-2530

TELEFRAX: (212) 840-3333

TELEFRAX: (212) 840-3133

TELEFRAX: (212) 840-3133

TELERX: 425066CURTMS

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.3%; Score 26.6; DB 1;
51.2%; Pred. No. 13;
tive 0; Mismatches 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Curtis, Morris & Safford
STREET: 530 Fifth Avenue
CITY: New York
                                                                                                                                                                                                                                                                                                                             US-08-184-009-215/c
; Sequence 215, Application US/08184009
; Patent No. 5833975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 1182 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 51.23
Conservative 62; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: cDNA
US-08-184-009-215
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                                                                           APPLICANT: Paoletti, Enzo
APPLICANT: Tartaglia, James
APPLICANT: Tortaglia, James
APPLICANT: COX, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/458,356
FILING DATE: 02-JUN-1995
CLASSIFICATION: 42-4
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/184,009
FILING DATE: 19-JAN-1994
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.3%; Score 26.6; DE 51.2%; Pred. No. 13; Live 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 454310-2530
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                     Sequence 215, Application US/08458356 Patent No. 5942235 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
                                                                                                                                                                                                                      : 530 Fifth Avenue
New York
                                                                                                                                                                                                                                                                                                                                                      Floppy disk
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TELEX: 42506CURTWG
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 1182 base pairs
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Best Local Similarity 51.2%
Matches 62% Conservative
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US-08-458-356-215
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

OM nucleic - nucleic search, using sw model

September 28, 2004, 12:07:16 ; Search time 15 Seconds (without alignments) 2.663 Million cell updates/sec Run on:

SEQ58-252-1430 Perfect acore:

l atggaggagccgcagtcaga......cagaagggcctgactcagac 1179 Sequence:

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10 segs, 16943 residues Searched:

Total number of hits satisfying chosen parameters:

20

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

\*: qpsbu Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARTES

		ф			SOUTHERES	
Result No.	Score	Query Match	Query Match Length DB	DB	ID	Description
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1	1179	100.0	1182	Н	AA067884	!
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c 12	26.6	2.3	1182		AA208529	Human ps3 DNA. Sy
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2 16	26.6			4 ,	0000000	He/ps3 (wildtype)
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)	•	7	400	4	ADD93286	p53-Yak3 fusion pr

ALIGNMENTS

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AA067884 standard; DNA; 1182 BP. (revised)
(first entry) 25-MAR-2003 23-MAR-1995 AAQ67884; AAQ67884 ID AAQ6 XX AC AAQ6 XX DT 25-h RESULT 1

Polymerase chain reaction; primer; amplify; NYVAC; ALVAC; recombinant; muxine; interleukin-2; IL-2; pRW825; pmut-1; pBS-SK; pMM151; TK vector; plasmid; vaccinia; H6 promoter; amplify; primer; antigenic response; polymerase chain reaction; poxvirus; pSb542; immunological response; pathogen; human; interferon; IFN; ss.

Synthetic.

WO9416716-A1.

04-AUG-1994.

21-JAN-1994;

94WO-US000888

93US-00007115. 19-JAN-1994; 21-JAN-1993;

(VIRO-) VIROGENETICS CORP.

Š Tartaglia J, Paoletti E,

ΜĬ,

WPI; 1994-263767/32

- comprises DNA Attenuated recombinant virus used for cancer therapy encoding cytokine and/or tumour associated antigen.

Example 32; Fig 39; 232pp; English.

This sequence represents the wildtype human p53 gene from the translation initiation codon to the stop codon. This sequence was used in the construction of an ALVAC-based recombinant virus containing a mutant form of the human p53 gene. The mutant form has a G-AB substitution at position plasmid pWMI10 (see also AAQ67864) contains the vaccinia H6 promoter and p53 gene was obtained from plasmid CA22A and cloned into pWMI10 to generate pWMI43. Recombination between pWMI43 and ALVAC rescuing virus produced recombinant virus vCP270, which contains the vaccinia H6 promoted mutated human p53 in the C5 locus. The resulting virus promoted mutated human p53 in the C5 locus. The resulting virus may be used in a composition for inducing an antigenic or immunological to correct PN field.) 

Sequence 1182 BP; 276 A; 365 C; 307 G; 234 T; 0 U; 0 Other;

Query Match

0 120 61 GACCTAIGGAAACTACTICCIGAAAACAACGITCIGICCCCIIGCCGICCCAAGCAAIG 120 9 61 GACCTATGGAAACTACTTCCTGAAAACAACGTTCTGTCCCCTTGCCGTCCCAAGCAATG 1 ATGGAGGAGCCGCAGTCAGATCCTAGCGTCGAGCCCCCTCTGAGTCAGGAAACATTTTCA Gaps ·, 100.0%; Score 1179; DB 1; Length 1182; 100.0%; Pred. No. 0.00058; Indels 0; Mismatches 0; Indels 0; Indels Best Local Similarity 100. Matches 1179; Conservative 임 8 ò g

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Attenuated recombinant virus; cytokine; tumour associated antigen; NYVAC recombinant virus; ALVAC recombinant virus; gene therapy; rabies; cancer; tumour necrosis factor; nuclear phosphoprotein; p53; IL-2; GMCSF;
                                                                                                                                Human p53 gene
AAZOB529
IID AAZO
XX AC AAZO
XX I9-C
XX DE Huma
XX Atte
XW Atte
KW NYVE
KW CANC
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BP.

entry)

19-OCT-1999 AAZ08529;

AAZ08529 standard; DNA; 1182

RESULT

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GACCTAIGGAAACTACTICCTGAAAACAACGIICTGICCCCCTIGCCGICCCAAGCAAIG 120

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The present invention describes a recombinant poxvirus (I), comprising exogenous DNA encoding an antigenic determinant of a pathogen which is exogenous DNA encoding an antigenic determinant of a pathogen which is then expressed in vivo in infected host calls after administration to a pathogen therefore induces an immunological response. (I) may be used to vaccinate patients against a wide range of diseases and disorders depending on the type of antigen encoded by the exogenous DNA. (I) may be used to vaccinate against diseases such as rabies, influenza and Newcatle Disease. It is particularly useful for immunising against cancers. The poxvirus (I) also provides a means of manipulating cancers. The poxvirus (I) also have enhanced safety compared to mattenuated viruses (attenuation reduces the virulence of the viruses) unattenuated viruses (attenuation reduces the virulence of the viruses) and known recombinant poxvirus vaccines. This increased level of safety reduces the possibility of a 'runaway' infection in the host and reduces the chance of transmission from vaccinated to unvaccinated individuals and contamination of the environment. The present sequence represents a constant of the environment. The present sequence represents a constant of the environment.
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interleukin; interferon; IFN-gamma; IL-4; melanoma associated antigen; carcinoembryonic antigen; immunisation; antigenic; poxvirus; influenza; immunological response; immunotherapy; vaccine; Newcastle Disease; ss.
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The present invention relates to a chimera gene of p53 family encoding a transcription activating region, a DNA binding region, and an oligomer formation region of different p53 family proteins. The chimeric gene can be used for gene therapy of p53 variant human tumours, and analysis of the function of the p53 family gene. The present sequence was used in the gene of the p53 family, useful for gene therapy, and treatment comprises a transcription activating region and a DNA binding 121 GATGATTTGATGCTGTCCCCGGACGATGTTGAACAATGGTTCACTGAAGACCCAGGTCCA GACCTATGGAAACTACTTCCTGAAAACAACGTTCTGTCCCCCTTGCCGTCCCAAGCAATG 61 GACCIALGRAAACIACITCCIGAAAACAACGTICTGICCCCTTGCCGTCCCAAGGAATG GATGATTIGATGCTGTCCCCGGACGATATTGAACAATGGTTCACTGAAGACCCAGGTCCA 1 ATGGAGGAGCCGCAGTCAGATCCTAGCGTCGAGCCCCTCTGAGTCAGGAAACATTTTCA AAAACCTACCAGGGAGCTACGGTTTCGTCTGGGCTTCTTGCATTCTGGGACAGCCAAG GATGAAGCTCCCAGAATGCCAGAGGCTGCTCCCCGGCGTGGCCCCTGCACCAGCAGCTCCT ACACCGGCGGCCCTGCACCAGCCCCTCCTGGCCCCTGTCATCTTCTGTCCCTTCCCAG 241 ACACCGGCCCCTGCACCCCTCCTGGCCCCTGTCATCTTCTGTCCTTCCCAG Gaps 0; 100.0%; Score 1179; DB 1; Length 1182; 100.0%; Pred. No. 0.00058; Live 0; Mismatches 0; Indels 0; Sequence 1182 BP; 276 A; 365 C; 307 G; 234 T; 0 U; 0 Other; AAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC 1179 gene therapy; p53; tumour; ss. Disclosure, Fig 2; 57pp; Japanese. CDNA; 1182 99JP-00139034. Conservative ဗ (IKAW/) IKAWA H. (SAKA ) OTSUKA PHARM WPI; 2001-268293/28. P-PSDB; AAB84836. p53 coding sequence standard; Best Local Similarity Matches 1179; Conser present invention JP2000354488-A Unidentified 09-APR-1999; 09-APR-1999; Cytostatic; 03-AUG-2001 AAH19387 AAH19387 Chimera c 61 Query Match 121 181 301 region, RESULT 3 AAH19387 엄 g ò 셤 qq ò ð 셤 ò a à d

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Attenuated recombinant virus; cytokine; tumour associated antigen; NYVAC recombinant virus; ALVAC recombinant virus; gene therapy; rabies; cancer; tumour necrosis factor; nuclear phosphoprotein; p53; IL-2; GMCSF; interleukin; interferon; IFN-gamma; IL-4; melanoma associated antigen; carcinoembryonic antigen; immunisation; antigenic; poxvirus; influenza; H6/p53 (wildtype) expression cassette and flanking regions from vCP207. ВР. AAZ08435 standard; DNA; 1484 19-OCT-1999 (first entry) AAZ08435; AAZ08435 RESULT 

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Disease;
immunological response; immunotherapy; vaccine; Newcastle
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88US-00234390.
90US-00537882.
91US-00637890.
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82US-00446824.
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                   Homo sapiens.
Vaccinia virus
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Recombinant poxviruses comprising exogenous DNA encoding antigenic determinants useful in immunotherapy to immunize against cancers and determinants useful in immunotherapy to immunize against cancer other diseases such as influenza, Newcastle Disease and rables.

Example 15; Fig 18; 163pp; English.

The present invention describes a recombinant poxvirus (I), comprising coordinate and therefore indicated host cells after administration to a patient and therefore induces an immunological response. (I) may be used to vaccinate patients against a wide range of diseases and disorders consider to vaccinate patients against a wide range of diseases and disorders consect to vaccinate against diseases buch as rabies, influenza and consect to vaccinate against diseases buch as rabies, influenza and consect. The poxvirus (I) also provides a means of manipulating cancers. The poxvirus (I) also have enhanced safety compared to modalities for cancer. (I) also have enhanced safety compared to concernate the poxvirus vaccines. This increased level of safety cand known recombinant poxvirus vaccines. This increased level of safety concernate the possibility of a 'runaway' infection in the host and reduces the chance of transmission from vaccinated to unvaccinated individuals containation of the environment. The present sequence represents \$6.00 (wildtype) expression cassette and flanking regions from vCP207 d in the exemplification of the present invention H6/p53 (wildtype) used in the exempl

Gaps . 0 100.0%; Score 1179; DB 1; Length 1484; 100.0%; Pred. No. 0.00047; ive 0; Mismatches 0; Indels 0; Sequence 1484 BP; 367 A; 416 C; 371 G; 330 T; 0 U; 0 Other; Similarity Query Match Best Local S

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Attenuated recombinant virus, cytokine, tumour associated antigen, NYVAC recombinant virus; dene therapy; rabies; cancer; tumour necrosis factor; nuclear phosphoprotein; p53; IL-2; GWCSF; interleukin; interferon; IEN-2gamma; IL-4; melanoma associated antigen; carcinoembryonic antigen; immunisation; antigenic; poxvirus; influenza; immunological response; immunotherapy; vaccine; Newcastle Disease; ss.
                                                (wildtype) expression cassette and flanking regions from vP1101
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Vaccinia virus.
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16-DEC-1991;
03-MAR-1992;
06-MAR-1992;
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INC. RES (HEAL-) HEALTH

Paoletti

WPI; 1999-493494/41.

Recombinant poxviruses comprising exogenous DNA encoding antigenic determinants useful in immunotherapy to immunize against cancers and other diseases such as influenza, Newcastle Disease and rabies.

Example 15; Fig 17; 163pp; English.

(I) may be be used The present invention describes a recombinant poxvirus (I), comprising exogenous DNA encoding an antigenic determinant of a pathogen which is then expressed in vivo in infected host cells after administration to a patient and therefore induces an immunological response. (I) may be use to vaccinate patients against a wide range of diseases and disorders depending on the type of antigen encoded by the exogenous DNA. (I) may be used to vaccinate against diseases such as rables, influenza and Newcastle Disease. It is particularly useful for immunising against cancers. The poxvirus (I) also provides a means of manipulating lymphocytes and tumour cells for use in cell-based immunotherapeutic

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modalities for cancer. (I) also have enhanced safety compared to unattenuated viruses (attenuation reduces the virulence of the viruses) and known recombinant poxvirus vaccines. This increased level of safety reduces the possibility of a 'runaway' infection in the host and reduces the chance of transmission from vaccinated to unvaccinated individuals and contamination of the environment. The present sequence represents a H6/p53 (wildtype) expression cassette and flanking regions from vP1101 used in the exemplification of the present invention
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iive 0; Mismatches 0;
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Best Local Similarity 100.
Matches 1179; Conservative
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This sequence encodes a p53-Chk1(1-270) fusion protein. This sequence may be used in the method of the invention for identifying a modulator of a specific serine/threonine kinase. The method comprises expressing a fusion protein between a substrate and a serine/threonine kinase within a cell, and determining the level of phosphorylation of the substrate in the presence of a candidate modulator. The method is useful for screening modulators of serine/threonine kinases. Serine/threonine kinases are
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                                                                                                                                                           GGGAGCAGGCTCACTCCAGCCACCTCAAAAAGGGTCAGTCTACCTCCCGCCAT
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                                   AAACCACTGGATGAGAATATTTCACCTTCAGATCCGTGGGCGTGAGCGCTTCGAGATG
                                                                                     CCAGGGAGCACTAAGCGAGCACTGCCCAACAACACCAGCTCCTCCCCCAGCCAAAGAAG
                                                                                                                                                                                                                                                                                                                                                                            ss; gene; p53-Chk1(1-270); fusion protein; modulation; serine/threonine kinase; phosphorylation; drug design; cancer.
                                                                                                                                                                                                                       AAAAAACTCATGTTCAAGACAGAGGGCCTGACTCAGAC 1179
                                                                                                                                                                                                                                                                                                                                                          coding sequence
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1. .2061
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                  Sequence 2061 BP; 552 A; 521 C; 531 G; 457 T; 0 U; 0 Other;
                                                                Indels
                                         DB 1;
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TTCCGAGAGCTGAATGAGGCCTTGGAACTCAAGGATGCCCAGGCTGGGAAGGAGCCAGGG
                                                               TTCCGAGAGCTGAATGAGGCCTTGGAACTCAAGGATGCCCAGGCTGGGAAGGAGCCAGGG
                                                                                              GGGAGCAGGGCTCACTCCAGCCACCTGAAGTCCAAAAAGGGTCAGTCTACCTCCCGCCAT
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                                                                                                                                                                                                                                                                                                                                                               ss; gene; p53-SGK(60-431); fusion protein; modulation; serine/threonine kinase; phosphorylation; drug design; cancer.
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                                                                                                                                                    AAAAAACTCATGTTCAAGACAGAGGGCCTGACTCAGAC 1179
                                                                                                                                                                       p53-SGK(60-431) fusion protein coding sequence.
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1. .2367
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100.0%; Score 1179; DB 1; Length 2367; 100.0%; Pred. No. 0.00029; 11ve 0; Mismatches 0; Indels 0;

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GGGAGCAGGCTCACTCCAGCCACCTGAAGTCCAAAAAAGGGTCAGTCTACCTCCCGCCAT
                                                                     AAAAAACTCATGTTCAAGACAGAGGGCCTGACTCAGAC 1179
                                                                                            coding sequence
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This sequence encodes a p53-Myt1(1-378) fusion protein. This sequence may be used in the method of the invention for identifying a modulator of a specific serine/threonine kinase. The method comprises expressing a fusion protein between a substrate and a serine/threonine kinase within a cell, and determining the level of phosphorylation of the substrate in the presence of a candidate modulator. The method is useful for screening modulators of serine/threonine kinases. Serine/threonine kinases are useful as targets for drug design, e.g., in the treatment of cancer.
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ss; gene; p53-Myt1(1-378); fusion protein; modulation; serine/threonine kinase; phosphorylation; drug design;
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                                                                                           p53-Yak3 fusion protein coding sequence.
                                                                                                                      ss; gene; p53-Yak3; fusion protein;
phosphorylation; drug design; cancer
                                                                                                                                                                                                                                      /product= "p53-Yak3"
                                                                                                                                                                                            Location/Qualifiers
1. .2406
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 7; Page 13; 38pp; English.
             ADD93286 standard; cDNA; 2406
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                                                                    (first entry)
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GENBANK; AF186773.
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Db ò ss.

cancer; breast cancer diagnosis; breast cancer antigen; gene;

15-MAY-2002; 2002US-00146473

US2003108888-A1

12-JUN-2003

Homo sapiens

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ADC35154 standard; cDNA; 1182

RESULT 10 ADC35154 ADC35154;

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                                                                                                                                                                                                                                                                                                    Diagnosing breast cancer in subject by obtaining biological sample from subject, contacting sample with breast cancer-associated polypeptides, determining specific binding between polypeptides and agents in sample.
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Pred. No. 0.00059;
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Polymerase chain reaction; primer; amplify; NYVAC; ALVAC; recombinant; murine; interleukin-2; IL-2; pRW825; pmut-1; pBS-SK; pMM151; TK vector; plasmid; vaccinia; H6 promoter; amplify; primer; antigenic response; polymerase chain reaction; poxvirus; pSD542; immunological response; pathogen; human; interferon; IFN; ss.
                         AAQ67884 standard; DNA; 1182
                                                                                                (first entry)
                                                                                  (revised)
                                                                               25-MAR-2003
23-MAR-1995
                                                                                                                        Human p53
                                                       AAQ67884;
RESULT 11
AAQ67884/c
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Synthetic

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This sequence represents the wildtype human p53 gene from the translation initiation codon to the stop codon. This sequence was used in the construction of an ALVAC-based recombinant virus containing a mutant form 524, changing an ALVAC-based recombinant virus containing a mutant form 524, changing an Arg residue at position 175 to a His residue. The plannid pwml10 (see also AAQ67864) contains the vaccinia H6 promoter and the wild type human p53 gene in the ALVAC 5 insertion site. The mutant generate pwml43. Recombination between pWM143 and ALVAC rescuing virus produced recombinant virus vCP270, which contains the vaccinia H6 promoted mutated human p53 in the C5 locus. The resulting virus promoted mutated human p53 in the C5 locus. The resulting virus may be used in a composition for inducing an antigenic or immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        234 AGCICCTACACCGGCGCCCCTGCACCAGCCCCTCCTGGCCCCTGTCATCTTCTGTCCC 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    174 AGGICCAGAIGAAGCICCCAGAAIGCCAGAGGCIGCICCCCGCGIGGCCCTGCACCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             294 AGGGACAGAAGATGACAGGGGCCAGGAGGGGCTGGTGCAGGGGCCGCCGCTGTAGGAGC
                                                                                                                                                                                                                - comprises DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               59; Indels
                                                                                                                                                                                                      Attenuated recombinant virus used for cancer therapy
                                                                                                                                                                                                                       encoding cytokine and/or tumour associated antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ch 2.3%; Score 26.6; I Similarity 51.2%; Pred. No. 12; 62; Conservative 0; Mismatches
                                                                                                                                                                                                                                              Example 32; Fig 39; 232pp; English.
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                                                                                                                                                      Cox
                                                                                 93US-00007115
94US-00184009
                                                        94WO-US000888
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                                                                                                                       (VIRO-) VIROGENETICS CORP.
                                                                                                                                                    Tartaglia J,
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                                                                                                                                                                             WPI; 1994-263767/32.
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WO9416716-A1
                                                      21-JAN-1994;
                                                                                             19-JAN-1994;
                           04-AUG-1994
                                                                                                                                                    Paoletti E,
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Best Local S:
Matches 62
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235

Attenuated recombinant virus; cytokine; tumour associated antigen; WYVAC recombinant virus; ALVAC recombinant virus; gene therapy; rabies; cancer; tumour necrosis factor; muchar phosphoprotein; p53; IL-2; GMCSF; interleukin; interferon; IEN-2gamma; IL-4; melanoma associated antigen; carcinoembryonic antigen; immunisation; antigent; poxvirus; influenza; immunological response; immunotherapy; vaccine; Newcastle Disease; ss.

Homo sapiens

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KX BX BX SX SX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a recombinant poxvirus (1), comprising exogenous DNA encoding an antigenic determinant of a pathogen which is chen expressed in vivo in infected host cells after administration to a patient and therefore induces a mimunological response. (1) may be used to vaccinate patients against a wide range of diseases and disorders depending on the type of antigen encoded by the exogenous DNA. (1) may be used to vaccinate against diseases such as rabies, influenza and Newcastle Disease. It is particularly useful for immunising against cancers. The poxvirus (1) also provides a means of manipularing cancers. The poxvirus (1) also have enhanced safety compared to modalities for cancer. (1) also have enhanced safety compared to unattenuated viruses (attenuation reduces the virulence of the viruses) reduces the possibility of a 'runaway' infection in the host and reduces the chance of transmission from the converse energy and converse reduces the chance of transmission from the converse represented the chance of transmission from the converse represented the converse that the converse reduces the converse reduces the chance of transmission from the converse represented the converse reduces the converse of transmission from the converse reduces the converse reduces the converse reduces the converse reduces the converse the converse reduces the converse reduces the converse the converse reduces the converse reduces the converse reduces the converse the converse reduces the converse reduces the converse 
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human p53 gene used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      234 AGCTCCTACACCGGCGGCCCCTGCACCAGCCCCTCTGGCCCCTGTCATCTTGTCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recombinant poxviruses comprising exogenous DNA encoding antigenic determinants useful in immunotherapy to immunize against cancers and other diseases such as influenza, Newcastle Disease and rabies.
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51.2%; Pred. No. 12;
tive 0; Mismatches
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                                                                                                                                81US-00334456.
82US-0046824.
84US-00902135.
87US-00090711.
87US-00110335.
88US-00186054.
88US-00234390.
90US-00537882.
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Best Local Similarity 51.4.
Best Local Similarity 51.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HEAL-) HEALTH RES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-493494/41.
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and contaminat
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20-OCT-1987;
25-APR-1988;
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14-JUN-1990;
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US5942235-A
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234 AGCICCIACACCGGCGGCCCCTGCACCACCCCCTCCTGGCCCCTGTCATCTTCTGTCCC 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimera gene of the p53 family, useful for gene therapy, and treatment of cancer, comprises a transcription activating region and a DNA binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a chimera gene of p53 family encoding a transcription activating region, a DNA binding region, and an oligomer formation region of different p53 family proteins. The chimeric gene can be used for gene therapy of p53 variant human tumours, and analysis of the function of the p53 family gene. The present sequence was used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           174 AGGTCCAGATGAAGCTCCCAGAATGCCAGAGGCTGCTCCCCGGGTGGCCCCTGCACCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast cancer; breast cancer diagnosis; breast cancer antigen; gene;
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Pred. No. 12;
0; Mismatches
                                                                                                                                                                                       Cytostatic; gene therapy; p53; tumour; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC35154 standard; cDNA; 1182 BP
               AAH19387 standard; cDNA; 1182 BP
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                                                                                                                                                                                                                                                                                                                                                                                                          99JP-00139034
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(SAKA ) OTSUKA PHARM CO LTD.
                                                                                                     (first entry)
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P-PSDB; AAB84836.
                                                                                                                                                p53 coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention
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                                                                                                                                                                                                                                                                                                                                                                   09-APR-1999;
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                                                                                                                                                                                                                                    Unidentified
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                                                               AAH19387;
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AAH19387/
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(LUDW-) LUDWIG INST CANCER RES
                            15-MAY-2002; 2002US-00146473.
                                     15-MAY-2001; 2001US-0291150P.
                                                      Scanlan MJ, Gout I,
                                                              WPI; 2003-829397/77
                                                                  P-PSDB; ADC35112.
            US2003108888-A1
    Homo sapiens
                    12-JUN-2003
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Chen Y;

Gure A,

old LJ,

Stockert E,

The invention describes a method of diagnosing breast cancer in subject comprising contacting biological sample from subject with at least two different breast cancer-associated polypeptides (I) encoded by nucleic acid molecules (II) comprising sequence chosen from 42 fully defined sequences as given in specification, determining specific binding between breast cancer. The method is useful for diagnostic for subject. The sample is blood, lymph node fluid or breast cancer in a This sequence encodes a breast cancer antigen. Diagnosing breast cancer in subject by obtaining biological sample from subject, contacting sample with breast cancer-associated polyperides, determining specific binding between polypeptides and agents in sample. Claim 1; SEQ ID NO 38; 173pp; English.

Sequence 1182 BP; 276 A; 364 C; 308 G; 234 T; 0 U; 0 Other;

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233
                                                                                           294 Addanchdahadhranchagacandanadagachdahadadagacacacadarhangahad 235
                                                                                                                        234 AGCTCCTACACCGGGGGGCCCCTGCACCAGCCCCTTCCTGGCCCCTGTCATCTTGTGTCCC 293
                                                                                                                                               234 TGCTGGTCCAGGGGCCACGGGGGAGCAGCCTCTGGGATTCTGGGAGCTTCATCTGGACC 175
                                                               174 AGGICCAGAIGAAGCICCCAGAATGCCAGAGGCTGCTCCCGCGTGGCCCCTGCACCAGC
                                               Gaps
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0
             DB 1; Length 1182;
                                           Indels
         Score 26.6; DB 1;
Pred. No. 12;
0; Mismatches 59;
      2.3%;
Query Match
Best Local Similarity 51.2'
Matches 62; Conservative
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RESULT 15 AAZ08415/c ID AAZ08435 standard, DNA, 1484 AAZ08435; 

19-OCT-1999 (first entry)

H6/p53 (wildtype) expression cassette and flanking regions from vCP207.

Attenuated recombinant virus; cytokine; tumour associated antigen; WVAC recombinant virus; ALVAC recombinant virus; gene therapy; rabies; cancer; tumour necrosis factor; nuclear phosphoprotein; p53; IL-2; GMCSF; intexleukin; interferon; IFN-gamma; IL-4; melanoma associated antigen; carcinoembryonic antigen; immunisation; antigenic; poxvirus; influenza; immunological response; immunotherapy; vaccine; Newcastle Disease; ss.

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Gaps

.; 0

59; Indels

0; Mismatches Score 26.6; D Pred. No. 9.9;

Conservative

2.3%;

Similarity

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Query Match Best Loca Matches

DB 1; Length 1484;

467

407

294 T 294

526 Addanchdhadhrdachagachachagadagachaghachagadachachaghadaga 234 AGCTCCTACACCGGCGCCCTGCACCAGCCCCCTCCTGGCCCCTGTCATCTTGTGCCC 466 Téchesieckagedecacedededeckectronecarronesascricatoridade

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> Homo sapiens. Vaccinia virus Synthetic.

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exogenous DNA encoding an antigenic determinant of a pathogen which is then expressed in vivo in infected host cells after administration to a pathogen which is then expressed in vivo in infected host cells after administration to a patient and therefore induces an immunological response. (I) may be used to vaccinate patients against a wide range of diseases and disorders depending on the type of antigen encoded by the exogenous DNA. (I) may be used to vaccinate against diseases such as rabies, influenza and loss of the use of the virulenza and cancers. The poxvirus (I) also provides a means of manipulating ancest. (I) also provides a means of manipulating modalities for cancer. (I) also have enhanced safety compared to unattenuated viruses (attenuation reduces the virulence of the viruses) and known recombinant poxvirus vaccines. This increased level of safety the chance of transmission from vaccinated to unvaccinated individuals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a recombinant poxvirus (I), comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and contamination of the environment. The present sequence represents (H6/p53 (Wildtype) expression cassette and flanking regions from vCP207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombinant poxviruses comprising exogenous DNA encoding antigenic determinants useful in immunotherapy to immunize against cancers and other diseases such as influenza, Newcastle Disease and rabies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             H6/p53 (wildtype) expression cassette and rianking reused in the exemplification of the present invention
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                                                                                                                                                                                    87US-00090711
87US-00110335
8BUS-00186054.
8BUS-00537882
90US-00537882.
91US-008057897.
92US-00847977.
92US-00847951.
92US-00847951.
92US-00847951.
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84US-00622135,
87US-00090209,
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94US-00184009.
94US-00228926.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HEAL-) HEALTH RES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-493494/41.
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Tue Sep 28 12:08:39 2004

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